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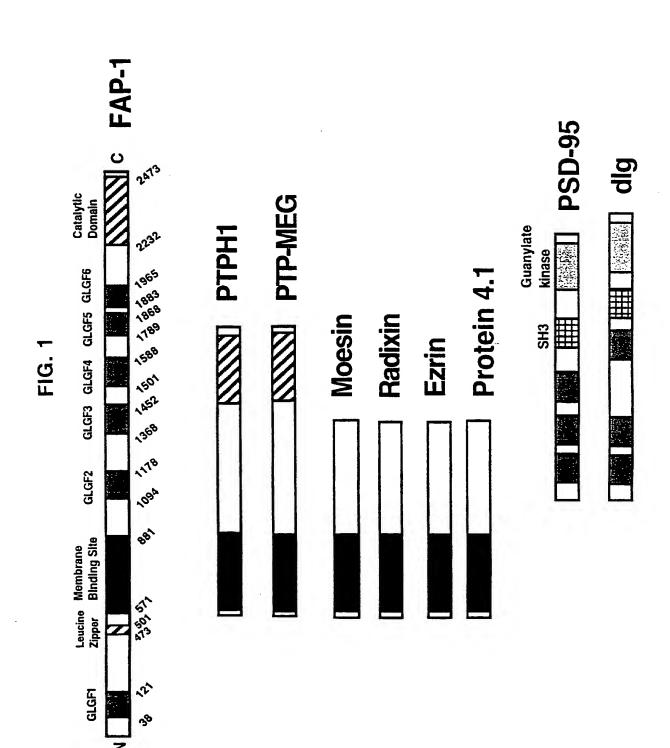
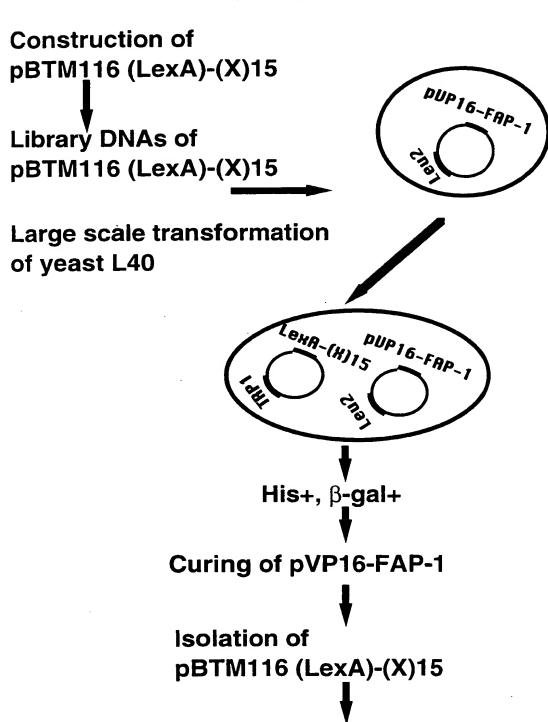


FIG. 2A



Analysis of DNA sequences

133

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EQSL

0-2

S

14-1

SNS

S

LA

57-5

S

6-3

V S O I

18-1

ESLV

16-13

S S L

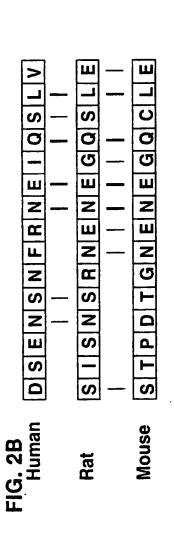
25-9

STV

Ш

72-1

SF



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S	
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FIG. 2D

FIG. 2C

IPPDSEDGNE	DSEMYNFRSQ	IDLASEFLFL	PPTCSQANSG	SDSNMNMN	QNFRTY	RET	RGF	E	
12-0	2-0	13-0	20-0	6-2	9-2	18-1	22-1	71-1	14-5
>	>	>	>	>	>	>	>	>	
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	ш	<u>a</u>	O	国		Z	z	A	1
		Н	Н			\mathbb{H}	H	H	H
5	G V S	٦ ع	O	S H	РР	SGV	RPV	M A Q	N E E
A	5	A	O			S	В	0	Z
H	H	H	H	H	H	H	H	H	H
		-	\vdash	\vdash				\vdash	
4	A	5	A	S	凹	口		X	H
CYA	ENA	W W G	EHA	N S S	GLR	GSD	DKK	T G K	ASR

Consensus: tS-X-V/L/I

FIG. 3A

Fas C terminal 15 a.a. peptide (μM)



200 -

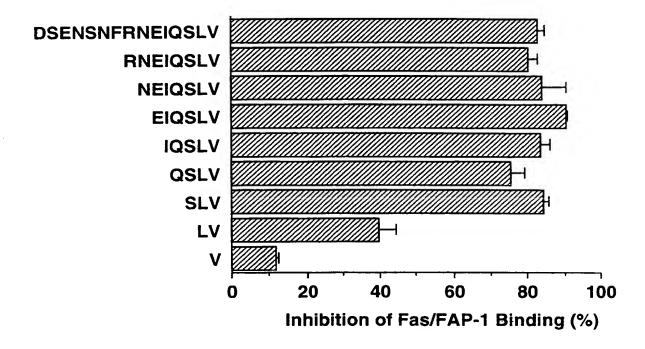
46 -

30 -21.5 -14.3 -

12345678910



FIG. 3B



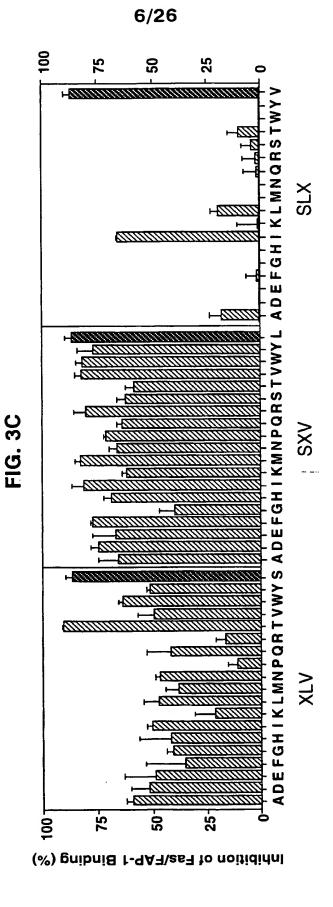
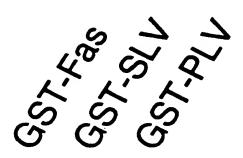


FIG. 4A

	VP16		VP16		
	FAP-1	Ras	FAP-1	Ras	
LexA					
Fas	700	000	6 0 0	金 卷 卷	
SLV	200	000		企业	
PLV			0 0 0	15 th 19	
SLY		8 6 6	国家企	e o c	
SLA			000		
	His	s +	Н	lis -	





250 -148 -

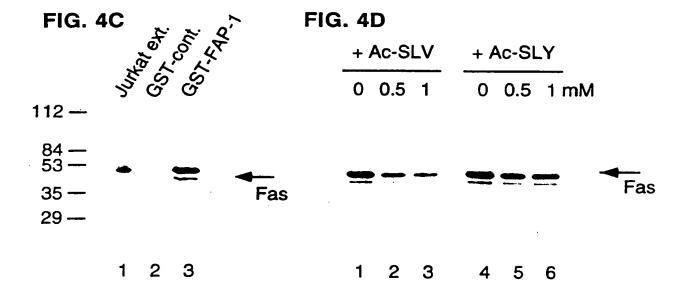
← FAP-1

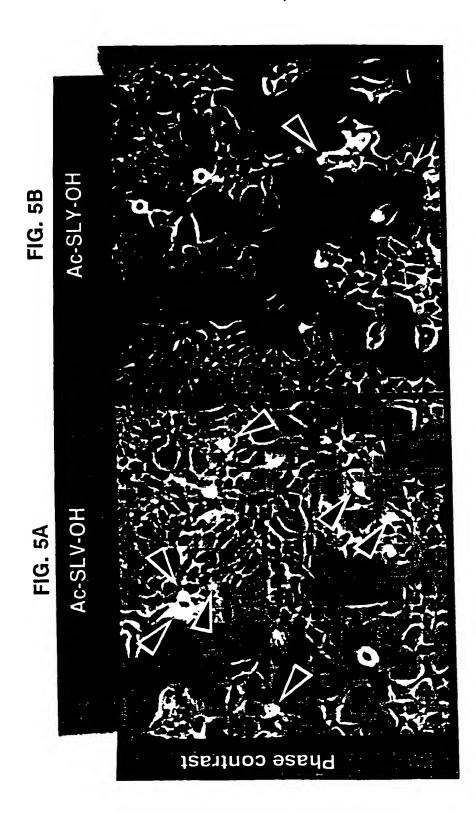
60 -

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1 2 3

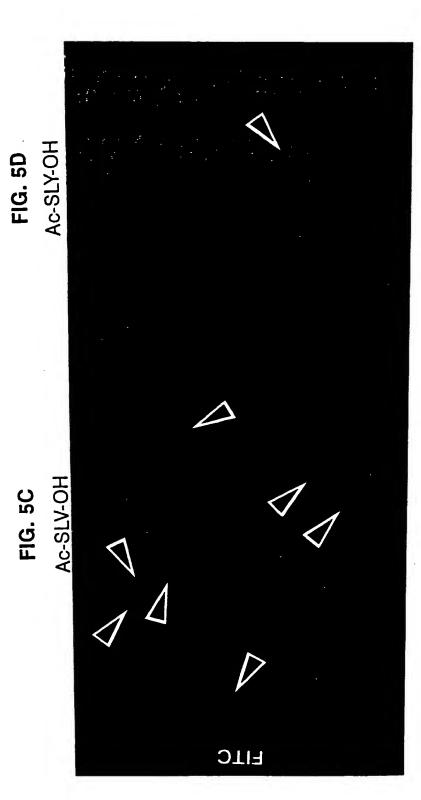




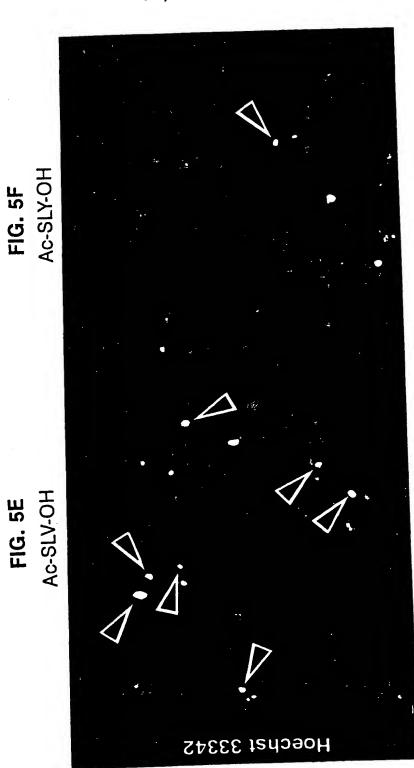


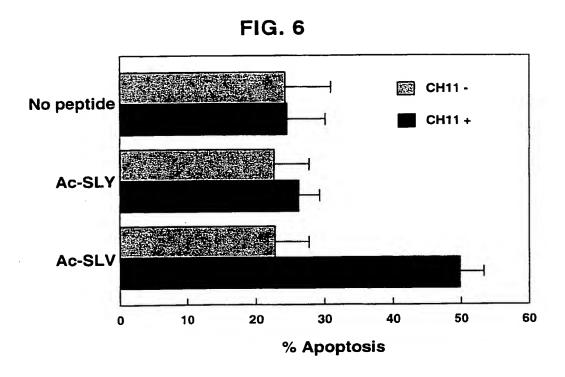






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vvttvmgssq egvaqpcgan edterglrec ngtpppegek adlveslcse ygyyddettg lngsagdtwr gecckacnlg eaddavcrca qdliastvag nkqgansrpv vdpclpctvc llaalrrigr pakreevekl acptglyths afkrwnsckg gldsmsapcv dgtysdeanh stdepeappe gdgglysslp atgdsatlda lgvslggake epckpctecv kgntvceecp ppegsdstap tqtasgqalk cpvrallasw avvvglvayi gsglvfscqd ipgrwitrst sqslhdqqph dgpr111111 vtfsdvvsat lipvycsila ehidsfthea qtvcepclds pvvtrgttdn mgagatgram trwadaecee rceacrvcea lhsdsgisvd hlagelgyqp statspv 181 241 361 121 301

Receptor

NGF

FIG. 7B

CD4 Receptor

kltgsgelww yagsgnltla vskrekavwv figlgiffcv ctasgkksig fhwknsngik vedqkeevql vsqlelqdsg iedsdtyice kniqggktls fsfplaftve hltlpgalpg slklenkeak vlggvaglll nfpliiknlk gkkgdtvelt vykkegegve psvqcrsprg klqmgkklpl stpvqpmali wgptspklml fqktc**spi** ekktcgcphr qlqknltcev dsrrslwdqg vsvkrvtqdp esnikvlptw ltlesppgss llvlglallp aatggkkvvl vlafqkassi kgpsklndra kkvefkidiv witfdlknke rmsqikrlls thllagaslt evnlvvmrat 11sdsgqv11 gaerasssks rcrhrrrgae lvfgltansd twtctvlqnq Inpeagmwdc ilgnggsflt leaktgklhg mnrgvpfrhl 301 181 241 361





FIG. 7C

Species	C-terminal sequences of NGFR (p75)	Binding activity of FAP-1
Human	SESTATSPV-COOH	+
Rat	SESTATSPV-COOH	+
Chicken	fSESTATSPV-COOH	+

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kklakaqceq wekelagire gpsspgrlts rttcsenela nlvaayekak selrselsgs nvvcqrkkss ksqndlltit agcsvqpwes esihidplsy issigvsssv slilgqfraa streagegay ssdrpvlgse erinsriehl gtererdlle elgrvitgle 111alaesed gttireedey ypnlaeersr gtrlqsvqat dgscggafav fvndlkrans aspalelael aavkitmlel kekkalelkl rphtnets. ysegcieaye naakallmkl rialleeens elkaqlylle dkpgkecada seirhqqsae elnkkidrig dadacsdins dyiqq1kadr divelnkriq klisktreess skirefevet drlrrrvrel etegvlgrdl natalrialq hetgvzmlkg lyshgsalse rahdcrktae slsstssgsk **vealeritks** elmamkeema hsaalaslkg ritelhsvia pengetmyta hodlaiktve atmnaireer **1plakiaerv** ftkedegr1k esquamver csniqeifqt msmlvgkyes gdenitomlk ldlenavlmg qerttlryee hiegittase skeeelnrtk elstssssnd kklkarvqel mdqdqtsvs1 sstasscdte veedkagrmr ndssaelsel tnrpinpstg aehlahsigd mnsgvamkyg shlmrehedv ghevnedsrs enesitamic gvgsepgdds Lssnahtstt dvkprgdsgr lvhiehlkse aeftnairre cslsvaevdr leecksnaer kkhqmk1kk; 181 241 301 421 481 481 541 661 601

dough that then the time then the lite he he he had then then that then

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pegdeegnme mlagqppfdg sptfcdhcgs sdfegfsyvn shctdfiwgf eklhvtvrđa gkvmladrkg drlyfvmeyv ldseghikia lkpsdkdrrl wwaygvllye talhecfatv lgcgpegerd dqlvianidq pownesttk arffkæptfc khkfkihtyg riylkaevad eegeyynvp1 flavlgkgsf yrdlkldnvm ktktirstin asgwykllng ldrvkltdfn ayqpygksvd glmtkhpakr cgmdhtekrg lfflhkrgii trgqpv1tpp kgpdtddprs lalldqppfl vhevkdhkfi Lskeavsick kgaenfdkff vfyaaeisig arkgalrqkn fvtiscpgad kgcvinvpsl popomeska fgvselmanp sedrkapsan vectimekry tpdyiapeii gpagnkvisp kkdvvigddd qvgkfkepqa dgvttrtfcg mehnvsypks trndfmgsls appfkpkvcg cfvvhkrche sdpyvklkl tasqdvanrf **ccdtcdmvh** 11yglihqgm knlipmdpng dweklenre1 gkagfacave nggdlmybiq ededelfqsi pathhpila madvfpgnds sveiwdwdrt teelyaikil dfgmckehmm 1rqkfekak1 661 361 421 481 541 601 181 241 301

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vntipalayk 1dryvaiqnp ddnfvligef escnedvíga seklfgrsih nrtalsced edgvnekv**gg v** enkkplqlil fsflpgssle vfkegsclla itrimavick dafnwtvdse lekklqmatn asimhlcais qhseeaskdn fsrylgcgyk pvfglqddsk gnilvimavs digtraklas 1fvvmwcpff dfnsgeants wlyldvlfst lgkeatlevs ackvlgivff ndcsmvalgk wplpsklcav tlfnktyrsa Inddtrlysn tavviiltia tisvgismpi lssavnplvy nskqdakttd vityfltiks tmgs1sneqk smitilygyr kaflkilavw lestrnslmg 1qekmwsa11 seqlqmqqkk repgaytgrr 11nvfvwigy mdilceents lspsclallh mligflumpv ibhsrfnsrt vsffipltim 121 181 241 301 361

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kalpnsgdet reskiyfrnp gdkteegver alltimfeam asmivtyflt dscrattlam tafikitww 1dtllltene atkevktlrk llvglfvmp1 vamldgsrkd igangynera fitnitivic eemkgiveeg gslaafftpl vdryjaikkp kerfgdfmlf detpcsspek stigestil waglqtesip flfllmwcpf yflmelavad mygspmrlrs stfvhvissn asimhleafs npnniccv1t lekklqyatn witvetvfqr raskvlgiví tlfnktfrda vknkppgr1t hgirnginpa kevatiened vesgvnplvy qstipehilq pikgietavd gntlvilavs wlfldvlfst 11sigiatov wplplvlcpa ilmvilptig Imrrtstigk lleifwigy ihalqkkayl maenskffkk malsyrvsel 7

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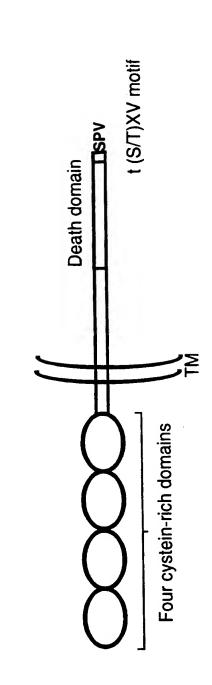
FIG. 7H

1 mazasydgii kqvealkmen snlrqeledn snhltklete asnmkevlkq lqgsiedeam 61 assggidlle rikelnidss nfpgvklrsk msirsygere gsvssrsgec spvpmgsfpr 121 rgfvngsres tgyleeleke rsllladidk eekekdwyya glamitkrid sipitenfsl 181 qtdmtrrqle yearqirvam eeqlgtcqdm ekraqrrlar iqqiekdilr irqllqsqa: 241 eaersequikh eigshdærg negggvæln mætagngggs tirmdnetas vissssthsa 301 przitshigt kvemvyslis migthdkódm srtliamses odscismrgs gcipiliqli 361 hgndkdsvli gnsrgskear arasaalhni ihsqpddkrg rreirvlhli eqiraycetc 421 wewgeahepg mdqddcnpmpa pvehqicpav cvlmklsfde ehrhamnelg glqaiaellq 481 vdcemygltn dhysitlrry agmaltnitf gdvankatic smkgcmraiv aqiksesedi 541 qqviasvlrn iswradvnsk ktirevgsvk almecalevk kestiksvis alwnisahct 601 erkadicavd galaflygtl tyrsqtntla iiesgggilr nyssliatne dhrqilrenn 661 clqtllqhlk shsltivsna cqtlwnlsar npkdqealwd mgavsmlknl ihskhkmiam 721 gsaaalrnim anrpakykda nimspgssip sihvrkqkal eaeldaqhis etfdnidnis 781 pkashrskqr hkqslygdyv fdtnrhddnr sdnfntgnmt vlspylnttv lpsssssrgs 841 ldssrsekdr slerergigl gnyhpatenp gtsskrglqi sttaaqiakv meevsaihts 901 qedrssgstt elhovtdern alrrssaaht hsntynftks ensnrtcsmp yakleykrss 961 ndelnevses dgygkrgqmk psiesysedd eskfcsyggy padlabkihs anhmddndge 1021 ldcpinyslk ysdeglnegr gspsqnerwa rpkhiledei kgsegrgsrn gsttypvyte 1081 stddkhlkfq phfgqqecvs pyrsrgangs etnrvgsnig inquvsqslc qeddyeddkp 1141 tnyserysee eqheeeerpt nysikyneek rhvdqpidys lkyatdipss qkqsfsfsks 1201 saggaskteh masssentst pasnakrong lhpssagsrs gopokaatok vasingetig 1261 tycvedtpic fsrcsslasl ssaedeigen gttgeadsan tlgiaeikek igtrsaedpv 1321 sevpaysonp rtkssrlogs slssesarhk avefssgaks paksgaotpk sppehyvoet 1381 plmferctsv ssldsfesrs iassvæsepc sæmvegiisp sdlpdspgæt mppsrektpp 1441 pppætaætkr evpknkapta ekresgpkæa avnaaværvæ vlpdadtilh fatestpæf 1501 scassisals ldepfickdv elrimppvce ndngmetese opkesnence keaektidse 1561 kdilddøddd dieileecii samptkserk akkpaqtask lpppvarkps qlpvykllps 1621 qarlqpqkhv sftpqddmpr vycvegtpin fstatsladl tiesppnela agegyrggaq 1681 sgefekrdti ptegrstdea gggktssvti pelddnkaee gdilaecins ampkgkshkp 1741 frykkimdov ogasasssap nknoldokkk ketspykpip onteyrtryr knadskonlin 1801 aeryfsdokd skkonlkons kdfodklipon edryrgsfaf dsphhytpie gtpycfsrod 1861 slssldfddd dydlsrekae lrkakenkes eakytshtel tsnggsankt gaiakgpinr 1921 gapkpilaka stfpasski pargastdek lanfaientp vofshnasis sisdidaenn 1981 nkenepiket eppdsogeps kpgasgyapk sfhvedtpvc fsrnsslssi sidseddllg 2041 ecissampkk kkpsrlkgdn ekhsprnmgg ilgedltldl kdigrpdseh glspdsenfd 2101 wkaigegans ivsslhqaaa aaclsrqass dsdsilslks gislgspfhl tpdqeekpft 2161 snkgprilkp gekstletkk ieseskgikg gkkvykslit gkvrsnseis ggmkqplqan 2221 mpsisrgrtm ihipgvinss sstspvskkg pplktpasks psegqtatts prgakpsvks 2281 elepvarqts qiggsskaps regerdstps rpaqqplsrp iqspgrnsis pgrngisppn 2341 kladlprtss patastkasg sgkmsytspg romsqqnltk qtglsknass iprsesaskg 2401 lnqmngnga nkkvelsrms stkssgsesd rserpvlvrq stfikeapsp tlrrkleesa 2461 sfeslspssr pasptrsqaq tpvlspslpd mslsthssvq aggwrklppn lsptieyndg 2521 rpakrhdiar shsespsrlp inrsgtwkre hskhssslpr vstwrrtgss ssilsasses 2581 sekaksedek hvnsisgtkq skenqvsakg twrkikenef sptnstsqtv ssgatngaes 2541 ktliygmapa vsktedvwvr iedopinnpr sgrsptgntp pvidsvseka npmikdskdn 2701 qakqnvgngs vpmrtvglen rlnsfiqvda pdqkgteikp gqnnpvpvse tnessivert 2761 pfsssssskh sspegtvaar vtpfnynpsp rkssadstsa rpsqiptpvn nntkkrdskt 2821 dstessgtqs pkrhsqsylv tsv

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p75NGFR

(Low-affinity nerve growth factor receptor)



first that then the could had

Fas

Fas

C-terminal amino acid sequence

NEIQSLV

STATSPV

STATSPV

PDZ domain t (S/T)-X-V |-COOH

interaction

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In vitro interaction of 35S-labeled FAP-1 with various receptors FAP-1 binds to the cytoplasmic region of p75NGFR.

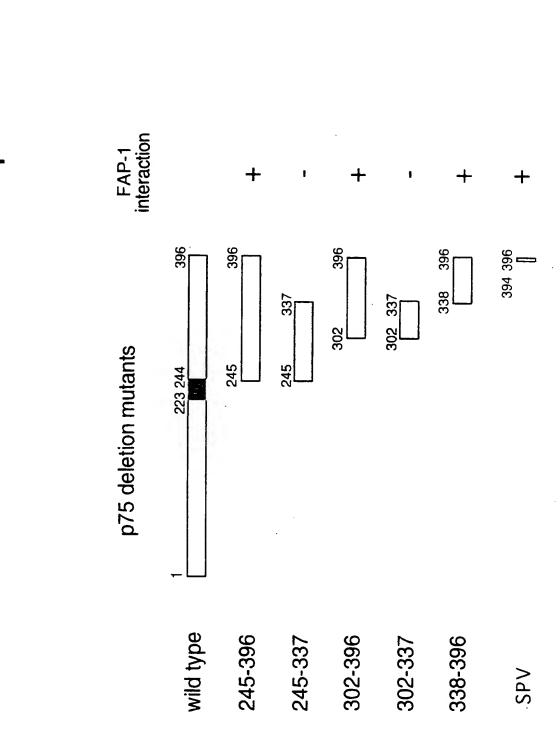
FIG. 10

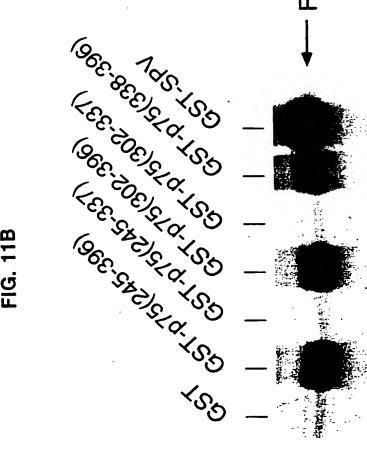
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1441 Hall 1850. Be will then 110 the the the Bolt Cine Weet Hall Cine 110.

FAP-1 binds to C-terminal three amino acids SPV of p75NGFR.

FIG. 11A





LexA-Lamin

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the first three to will hast

FAP-1 binds to p75NGFR C-terminal cytoplasmic region in yeast.

VP16-cRaf VP16-FAP-1 LexA-p75NGFR(338-396) LexA-p75NGFR(365-396) LexA-Ras^{V12} LexA-Fas